```
April 16, 2005, 05:19:07; Search time 34 Seconds (without alignments) 20.550 Million cell updates/sec
                                                                                                                                                                                                                                                    306091
                                                                                                                                                                                                                                                                                                                                                                  Pending_Patents_AA_New:*
: /cgn2_6/ptodata/2/paa/PCT_NEW_COMB.pep:*
:: /cgn2_6/ptcdata/2/paa/USO6_NEW_COMB.pep:*
:: /cgn2_6/ptc
          GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                        306091 seqs, 69870781 residues
                                                           OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                              Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                      BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                          Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                   US-09-018-194-4
57
1 CVGSNKGAIC 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. is the number c
score greater than or equand is derived by analysi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query
Match Length DE
                                                                                                                                   Title:
Perfect score:
Sequence:
                                                                                                                                                                                      Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
                                                                                                                                                                                                                                                                                                                                                               Database :
                                                                                                                                                                                                                           Searched:
                                                                                     Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Result
No.
```

Weger 1, 3. 09 1018 194 Page 1 59.10 4 Interf

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OM protein - protein search, using sw model

Run on:

April 16, 2005, 05:07:37; Search time 489 Seconds (without alignments) 23.886 Million cell updates/sec

US-09-018-194-4 57 1 CVGSNKGAIC 10 Title: Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

6959266 6959266 seqs, 1168006243 residues Searched:

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Pending Patents AA Main: * Database : ...

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Length	1123 1233 1233 1233 1233 1233 1233 1233		
Query Match	11000 1000 1000 1000 1000 1000 1000 10	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	
Score		,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	
Result No.	110 6 4 8 8 9 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		7 8 8 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9

RESULT 1 US-09-01	RESULT 1 US-09-018-194-4
. Se	Sequence 4; Application US/09018194
·-	GENERAL INFORMATION:
	APPLICANT: Gilchrest, Barbara A.
	APPLICANT: Yaar, Mina
٠.	APPLICANT: Eller, Mark
•	TITLE OF INVENTION: METHODS OF INDUCING HAIR GROWTH AND
	TITLE OF INVENTION: COLORATION
	NUMBER OF SEQUENCES: 10
	CORRESPONDENCE ADDRESS:
	ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
	STREET: Two Militia Drive
	CITY: Lexington
	STATE: Massachusetts
	COUNTRY: USA
	ZIP: 02173
	COMPUTER READABLE FORM:
	MEDIUM TYPE: Ploppy disk

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